

=====

Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=8; day=28; hr=9; min=5; sec=47; ms=172;]

=====

Application No: 10547842 Version No: 2.0

Input Set:

Output Set:

Started: 2008-07-25 10:52:04.840
Finished: 2008-07-25 10:52:07.749
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 909 ms
Total Warnings: 4
Total Errors: 0
No. of SeqIDs Defined: 5
Actual SeqID Count: 5

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)

SEQUENCE LISTING

<110> PINNA, LORENZO
 DONELLA-DEANA, ARIANNA
 MARIN, ORIANO
 MOLOGNI, LUCA
 GUNBY, ROSALIND
 GAMBACORTI PASSERINI, CARLO
 SCAPOZZA, LEONARDO

<120> ANAPLASTIC LYMPHOMA KINASE ASSAY, REAGENTS AND
 COMPOSITIONS THEREOF

<130> 2503-1169

<140> 10547842

<141> 2006-05-15

<150> PCT/EP2004/002185

<151> 2004-03-04

<150> EP 03005186.6

<151> 2003-03-07

<160> 5

<170> PatentIn Ver. 3.3

<210> 1

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 peptide

<400> 1

Ala	Arg	Asp	Ile	Tyr	Arg	Ala	Ser	Phe	Phe	Arg	Lys	Gly	Gly	Cys	Ala
1															
				5							10				15

Met	Leu	Pro	Val	Lys
				20

<210> 2

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 peptide

<400> 2

Ala Arg Asp Ile Tyr Arg Ala Ser Tyr Tyr Arg Lys Gly Gly Cys Ala
1 5 10 15

Met Leu Pro Val Lys
20

<210> 3

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
peptide

<400> 3

Ala Arg Asp Ile Phe Arg Ala Ser Tyr Phe Arg Lys Gly Gly Cys Ala
1 5 10 15

Met Leu Pro Val Lys
20

<210> 4

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
peptide

<400> 4

Ala Arg Asp Ile Phe Arg Ala Ser Phe Tyr Arg Lys Gly Gly Cys Ala
1 5 10 15

Met Leu Pro Val Lys
20

<210> 5

<211> 1620

<212> PRT

<213> Homo sapiens

<400> 5

Met Gly Ala Ile Gly Leu Leu Trp Leu Leu Pro Leu Leu Leu Ser Thr
1 5 10 15

Ala Ala Val Gly Ser Gly Met Gly Thr Gly Gln Arg Ala Gly Ser Pro
20 25 30

Ala Ala Gly Ser Pro Leu Gln Pro Arg Glu Pro Leu Ser Tyr Ser Arg
35 40 45

Leu	Gln	Arg	Lys	Ser	Leu	Ala	Val	Asp	Phe	Val	Val	Pro	Ser	Leu	Phe	50	55	60	
Arg	Val	Tyr	Ala	Arg	Asp	Leu	Leu	Leu	Pro	Pro	Ser	Ser	Ser	Glu	Leu	65	70	75	80
Lys	Ala	Gly	Arg	Pro	Glu	Ala	Arg	Gly	Ser	Leu	Ala	Leu	Asp	Cys	Ala	85	90	95	
Pro	Leu	Leu	Arg	Leu	Leu	Gly	Pro	Ala	Pro	Gly	Val	Ser	Trp	Thr	Ala	100	105	110	
Gly	Ser	Pro	Ala	Pro	Ala	Glu	Ala	Arg	Thr	Leu	Ser	Arg	Val	Leu	Lys	115	120	125	
Gly	Gly	Ser	Val	Arg	Lys	Leu	Arg	Arg	Ala	Lys	Gln	Leu	Val	Leu	Glu	130	135	140	
Leu	Gly	Glu	Glu	Ala	Ile	Leu	Glu	Gly	Cys	Val	Gly	Pro	Pro	Gly	Glu	145	150	155	160
Ala	Ala	Val	Gly	Leu	Leu	Gln	Phe	Asn	Leu	Ser	Glu	Leu	Phe	Ser	Trp	165	170	175	
Trp	Ile	Arg	Gln	Gly	Glu	Gly	Arg	Leu	Arg	Ile	Arg	Leu	Met	Pro	Glu	180	185	190	
Lys	Lys	Ala	Ser	Glu	Val	Gly	Arg	Glu	Gly	Arg	Leu	Ser	Ala	Ala	Ile	195	200	205	
Arg	Ala	Ser	Gln	Pro	Arg	Leu	Leu	Phe	Gln	Ile	Phe	Gly	Thr	Gly	His	210	215	220	
Ser	Ser	Leu	Glu	Ser	Pro	Thr	Asn	Met	Pro	Ser	Pro	Ser	Pro	Asp	Tyr	225	230	235	240
Phe	Thr	Trp	Asn	Leu	Thr	Trp	Ile	Met	Lys	Asp	Ser	Phe	Pro	Phe	Leu	245	250	255	
Ser	His	Arg	Ser	Arg	Tyr	Gly	Leu	Glu	Cys	Ser	Phe	Asp	Phe	Pro	Cys	260	265	270	
Glu	Leu	Glu	Tyr	Ser	Pro	Pro	Leu	His	Asp	Leu	Arg	Asn	Gln	Ser	Trp	275	280	285	
Ser	Trp	Arg	Arg	Ile	Pro	Ser	Glu	Glu	Ala	Ser	Gln	Met	Asp	Leu	Leu	290	295	300	
Asp	Gly	Pro	Gly	Ala	Glu	Arg	Ser	Lys	Glu	Met	Pro	Arg	Gly	Ser	Phe	305	310	315	320
Leu	Leu	Leu	Asn	Thr	Ser	Ala	Asp	Ser	Lys	His	Thr	Ile	Leu	Ser	Pro	325	330	335	
Trp	Met	Arg	Ser	Ser	Ser	Glu	His	Cys	Thr	Leu	Ala	Val	Ser	Val	His	340	345	350	

Arg	His	Leu	Gln	Pro	Ser	Gly	Arg	Tyr	Ile	Ala	Gln	Leu	Leu	Pro	His	355	360	365	
Asn	Glu	Ala	Ala	Arg	Glu	Ile	Leu	Leu	Met	Pro	Thr	Pro	Gly	Lys	His	370	375	380	
Gly	Trp	Thr	Val	Leu	Gln	Gly	Arg	Ile	Gly	Arg	Pro	Asp	Asn	Pro	Phe	385	390	395	400
Arg	Val	Ala	Leu	Glu	Tyr	Ile	Ser	Ser	Gly	Asn	Arg	Ser	Leu	Ser	Ala	405	410	415	
Val	Asp	Phe	Phe	Ala	Leu	Lys	Asn	Cys	Ser	Glu	Gly	Thr	Ser	Pro	Gly	420	425	430	
Ser	Lys	Met	Ala	Leu	Gln	Ser	Ser	Phe	Thr	Cys	Trp	Asn	Gly	Thr	Val	435	440	445	
Leu	Gln	Leu	Gly	Gln	Ala	Cys	Asp	Phe	His	Gln	Asp	Cys	Ala	Gln	Gly	450	455	460	
Glu	Asp	Glu	Ser	Gln	Met	Cys	Arg	Lys	Leu	Pro	Val	Gly	Phe	Tyr	Cys	465	470	475	480
Asn	Phe	Glu	Asp	Gly	Phe	Cys	Gly	Trp	Thr	Gln	Gly	Thr	Leu	Ser	Pro	485	490	495	
His	Thr	Pro	Gln	Trp	Gln	Val	Arg	Thr	Leu	Lys	Asp	Ala	Arg	Phe	Gln	500	505	510	
Asp	His	Gln	Asp	His	Ala	Leu	Leu	Leu	Ser	Thr	Thr	Asp	Val	Pro	Ala	515	520	525	
Ser	Glu	Ser	Ala	Thr	Val	Thr	Ser	Ala	Thr	Phe	Pro	Ala	Pro	Ile	Lys	530	535	540	
Ser	Ser	Pro	Cys	Glu	Leu	Arg	Met	Ser	Trp	Leu	Ile	Arg	Gly	Val	Leu	545	550	555	560
Arg	Gly	Asn	Val	Ser	Leu	Val	Leu	Val	Glu	Asn	Lys	Thr	Gly	Lys	Glu	565	570	575	
Gln	Gly	Arg	Met	Val	Trp	His	Val	Ala	Ala	Tyr	Glu	Gly	Leu	Ser	Leu	580	585	590	
Trp	Gln	Trp	Met	Val	Leu	Pro	Leu	Leu	Asp	Val	Ser	Asp	Arg	Phe	Trp	595	600	605	
Leu	Gln	Met	Val	Ala	Trp	Trp	Gly	Gln	Gly	Ser	Arg	Ala	Ile	Val	Ala	610	615	620	
Phe	Asp	Asn	Ile	Ser	Ile	Ser	Leu	Asp	Cys	Tyr	Leu	Thr	Ile	Ser	Gly	625	630	635	640
Glu	Asp	Lys	Ile	Leu	Gln	Asn	Thr	Ala	Pro	Lys	Ser	Arg	Asn	Leu	Phe	645	650	655	

Glu	Arg	Asn	Pro	Asn	Lys	Glu	Leu	Lys	Pro	Gly	Glu	Asn	Ser	Pro	Arg	
			660						665			670				
Gln	Thr	Pro	Ile	Phe	Asp	Pro	Thr	Val	His	Trp	Leu	Phe	Thr	Thr	Cys	
			675						680			685				
Gly	Ala	Ser	Gly	Pro	His	Gly	Pro	Thr	Gln	Ala	Gln	Cys	Asn	Asn	Ala	
			690						695			700				
Tyr	Gln	Asn	Ser	Asn	Leu	Ser	Val	Glu	Val	Gly	Ser	Glu	Gly	Pro	Leu	
705						710						715			720	
Lys	Gly	Ile	Gln	Ile	Trp	Lys	Val	Pro	Ala	Thr	Asp	Thr	Tyr	Ser	Ile	
			725						730						735	
Ser	Gly	Tyr	Gly	Ala	Ala	Gly	Gly	Lys	Gly	Gly	Lys	Asn	Thr	Met	Met	
			740						745			750				
Arg	Ser	His	Gly	Val	Ser	Val	Leu	Gly	Ile	Phe	Asn	Leu	Glu	Lys	Asp	
			755						760			765				
Asp	Met	Leu	Tyr	Ile	Leu	Val	Gly	Gln	Gln	Gly	Glu	Asp	Ala	Cys	Pro	
770						775						780				
Ser	Thr	Asn	Gln	Leu	Ile	Gln	Lys	Val	Cys	Ile	Gly	Glu	Asn	Asn	Val	
785						790						795			800	
Ile	Glu	Glu	Glu	Ile	Arg	Val	Asn	Arg	Ser	Val	His	Glu	Trp	Ala	Gly	
			805						810						815	
Gly	Gly	Gly	Gly	Gly	Gly	Gly	Ala	Thr	Tyr	Val	Phe	Lys	Met	Lys	Asp	
			820						825						830	
Gly	Val	Pro	Val	Pro	Leu	Ile	Ile	Ala	Ala	Gly	Gly	Gly	Gly	Arg	Ala	
			835						840			845				
Tyr	Gly	Ala	Lys	Thr	Asp	Thr	Phe	His	Pro	Glu	Arg	Leu	Glu	Asn	Asn	
850						855						860				
Ser	Ser	Val	Leu	Gly	Leu	Asn	Gly	Asn	Ser	Gly	Ala	Ala	Gly	Gly	Gly	
865						870						875			880	
Gly	Gly	Trp	Asn	Asp	Asn	Thr	Ser	Leu	Leu	Trp	Ala	Gly	Lys	Ser	Leu	
			885						890						895	
Gln	Glu	Gly	Ala	Thr	Gly	Gly	His	Ser	Cys	Pro	Gln	Ala	Met	Lys	Lys	
			900						905						910	
Trp	Gly	Trp	Glu	Thr	Arg	Gly	Gly	Phe	Gly	Gly	Gly	Gly	Gly	Gly	Cys	
915						920						925				
Ser	Ser	Gly	Gly	Gly	Gly	Gly	Gly	Tyr	Ile	Gly	Gly	Asn	Ala	Ala	Ser	
930						935						940				
Asn	Asn	Asp	Pro	Glu	Met	Asp	Gly	Glu	Asp	Gly	Val	Ser	Phe	Ile	Ser	
945						950						955			960	

Pro Leu Gly Ile Leu Tyr Thr Pro Ala Leu Lys Val Met Glu Gly His		
965	970	975
Gly Glu Val Asn Ile Lys His Tyr Leu Asn Cys Ser His Cys Glu Val		
980	985	990
Asp Glu Cys His Met Asp Pro Glu Ser His Lys Val Ile Cys Phe Cys		
995	1000	1005
Asp His Gly Thr Val Leu Ala Glu Asp Gly Val Ser Cys Ile Val Ser		
1010	1015	1020
Pro Thr Pro Glu Pro His Leu Pro Leu Ser Leu Ile Leu Ser Val Val		
1025	1030	1035
Thr Ser Ala Leu Val Ala Ala Leu Val Leu Ala Phe Ser Gly Ile Met		
1045	1050	1055
Ile Val Tyr Arg Arg Lys His Gln Glu Leu Gln Ala Met Gln Met Glu		
1060	1065	1070
Leu Gln Ser Pro Glu Tyr Lys Leu Ser Lys Leu Arg Thr Ser Thr Ile		
1075	1080	1085
Met Thr Asp Tyr Asn Pro Asn Tyr Cys Phe Ala Gly Lys Thr Ser Ser		
1090	1095	1100
Ile Ser Asp Leu Lys Glu Val Pro Arg Lys Asn Ile Thr Leu Ile Arg		
1105	1110	1115
Gly Leu Gly His Gly Ala Phe Gly Glu Val Tyr Glu Gly Gln Val Ser		
1125	1130	1135
Gly Met Pro Asn Asp Pro Ser Pro Leu Gln Val Ala Val Lys Thr Leu		
1140	1145	1150
Pro Glu Val Cys Ser Glu Gln Asp Glu Leu Asp Phe Leu Met Glu Ala		
1155	1160	1165
Leu Ile Ile Ser Lys Phe Asn His Gln Asn Ile Val Arg Cys Ile Gly		
1170	1175	1180
Val Ser Leu Gln Ser Leu Pro Arg Phe Ile Leu Leu Glu Leu Met Ala		
1185	1190	1195
Gly Gly Asp Leu Lys Ser Phe Leu Arg Glu Thr Arg Pro Arg Pro Ser		
1205	1210	1215
Gln Pro Ser Ser Leu Ala Met Leu Asp Leu Leu His Val Ala Arg Asp		
1220	1225	1230
Ile Ala Cys Gly Cys Gln Tyr Leu Glu Glu Asn His Phe Ile His Arg		
1235	1240	1245
Asp Ile Ala Ala Arg Asn Cys Leu Leu Thr Cys Pro Gly Pro Gly Arg		
1250	1255	1260

Val Ala Lys Ile Gly Asp Phe Gly Met Ala Arg Asp Ile Tyr Arg Ala			
1265	1270	1275	1280
Ser Tyr Tyr Arg Lys Gly Gly Cys Ala Met Leu Pro Val Lys Trp Met			
	1285	1290	1295
Pro Pro Glu Ala Phe Met Glu Gly Ile Phe Thr Ser Lys Thr Asp Thr			
	1300	1305	1310
Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe Ser Leu Gly Tyr Met			
	1315	1320	1325
Pro Tyr Pro Ser Lys Ser Asn Gln Glu Val Leu Glu Phe Val Thr Ser			
	1330	1335	1340
Gly Gly Arg Met Asp Pro Pro Lys Asn Cys Pro Gly Pro Val Tyr Arg			
1345	1350	1355	1360
Ile Met Thr Gln Cys Trp Gln His Gln Pro Glu Asp Arg Pro Asn Phe			
	1365	1370	1375
Ala Ile Ile Leu Glu Arg Ile Glu Tyr Cys Thr Gln Asp Pro Asp Val			
	1380	1385	1390
Ile Asn Thr Ala Leu Pro Ile Glu Tyr Gly Pro Leu Val Glu Glu Glu			
	1395	1400	1405
Glu Lys Val Pro Val Arg Pro Lys Asp Pro Glu Gly Val Pro Pro Leu			
	1410	1415	1420
Leu Val Ser Gln Gln Ala Lys Arg Glu Glu Glu Arg Ser Pro Ala Ala			
1425	1430	1435	1440
Pro Pro Pro Leu Pro Thr Thr Ser Ser Gly Lys Ala Ala Lys Lys Pro			
	1445	1450	1455
Thr Ala Ala Glu Val Ser Val Arg Val Pro Arg Gly Pro Ala Val Glu			
	1460	1465	1470
Gly Gly His Val Asn Met Ala Phe Ser Gln Ser Asn Pro Pro Ser Glu			
	1475	1480	1485
Leu His Lys Val His Gly Ser Arg Asn Lys Pro Thr Ser Leu Trp Asn			
	1490	1495	1500
Pro Thr Tyr Gly Ser Trp Phe Thr Glu Lys Pro Thr Lys Lys Asn Asn			
1505	1510	1515	1520
Pro Ile Ala Lys Lys Glu Pro His Asp Arg Gly Asn Leu Gly Leu Glu			
	1525	1530	1535
Gly Ser Cys Thr Val Pro Pro Asn Val Ala Thr Gly Arg Leu Pro Gly			
	1540	1545	1550
Ala Ser Leu Leu Leu Glu Pro Ser Ser Leu Thr Ala Asn Met Lys Glu			
	1555	1560	1565

Val Pro Leu Phe Arg Leu Arg His Phe Pro Cys Gly Asn Val Asn Tyr
1570 1575 1580

Gly Tyr Gln Gln Gln Gly Leu Pro Leu Glu Ala Ala Thr Ala Pro Gly
1585 1590 1595 1600

Ala Gly His Tyr Glu Asp Thr Ile Leu Lys Ser Lys Asn Ser Met Asn
1605 1610 1615

Gln Pro Gly Pro
1620